

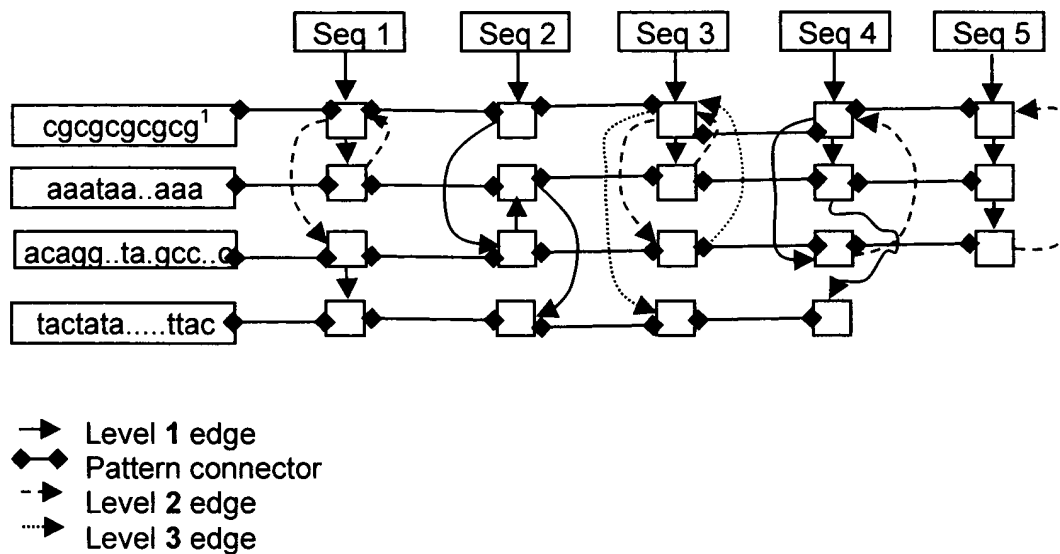
10/699,024

7-28-2006  
Replacement Sheet

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400

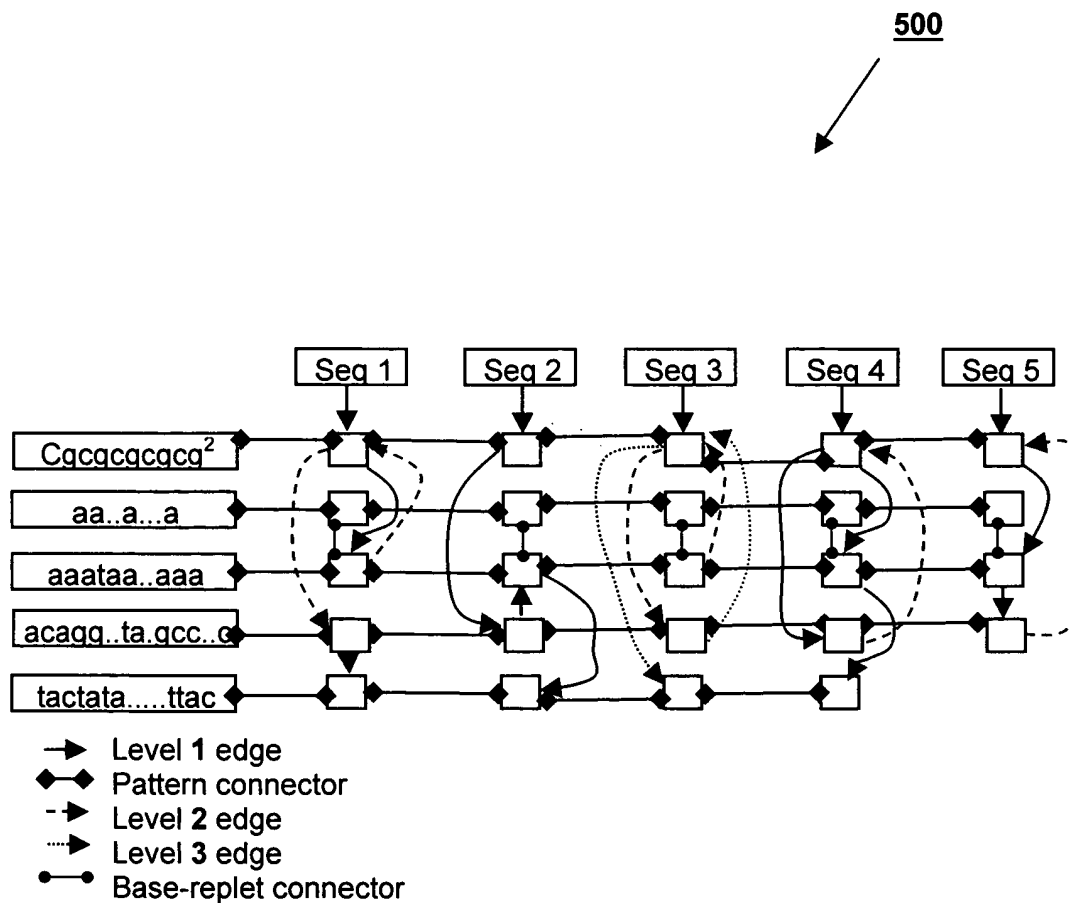


Footnote:  
1. sequence id 1 in accompanying sequence listing

FIG. 4

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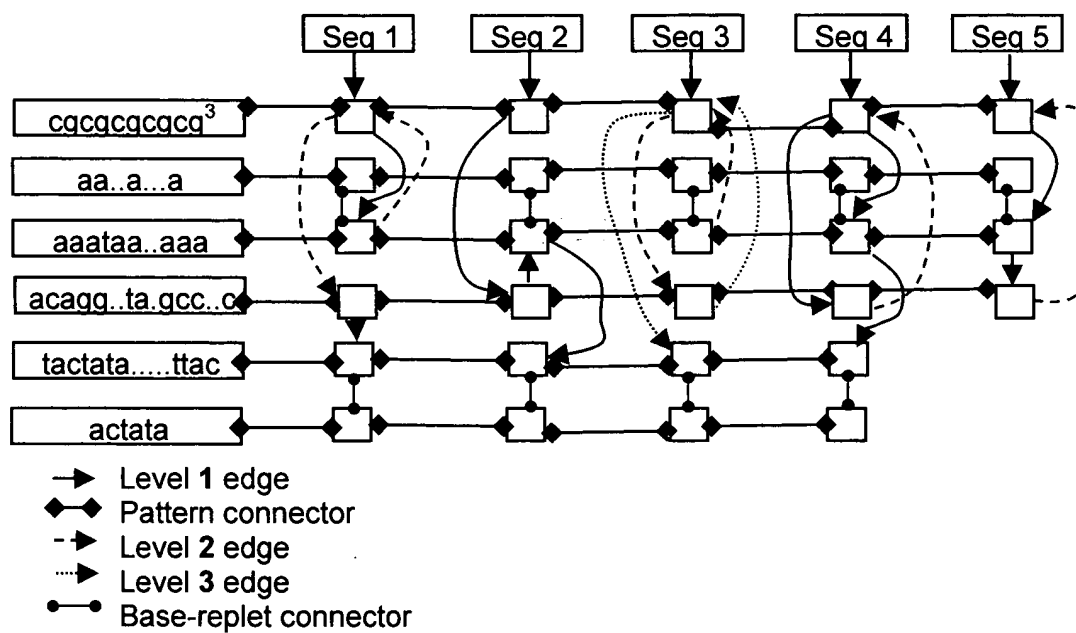


Footnote:  
2. sequence id 1 in accompanying sequence listing

FIG. 5

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Footnote:

3. sequence id 1 in accompanying sequence listing

FIG. 6

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Backbone = bseq 3: acttgatcggtagctagacggagaagctcccaaac

Base reptlets occurring in 3 are {cgcgcgcgcg<sup>1</sup>, aaataa..aaa, acagg..ta.gcc..c, tactata.....ttac}

Match-set of the base reptlets are provided below

**1: cgcgcgcgcg<sup>4</sup>**

```
{
Sequence-id = 3
Pattern-id = 1
Array of Matching-offsets <K,δ> = {18,39,83}
Array of Is-base-replet = {true, true, true}
Array of Pointer to Base-replet = {null, null, null}
Array of sequence-formation-edges = {2, 3, 4}
Pointer to next-pattern instance = {...}, Pointer to previous-pattern instance = {...}
}
```

**2: aaataa..aaa**

```
{
Sequence-id = 3
Pattern-id = 2
Array of Matching-offsets <K,δ> = {28}
Array of Is-base-replet = {true}
Array of Pointer to Base-replet = {null}
Array of sequence-formation-edges = {1}
Pointer to next-pattern instance = {...}, Pointer to previous-pattern instance = {...}
}
```

**3: acagg..ta.gcc..c**

```
{
Sequence-id = 3
Pattern-id = 3
Array of Matching-offsets <K,δ> = {49}
Array of Is-base-replet = {true}
Array of Pointer to Base-replet = {null}
Array of sequence-formation-edges = {1}
Pointer to next-pattern instance = {...}, Pointer to previous-pattern instance = {...}
}
```

**4: tactata.....ttac**

```
{
Sequence-id = 3
Pattern-id = 4
Array of Matching-offsets <K,δ> = {93}
Array of Is-base-replet = {true}
Array of Pointer to Base-replet = {null}
Array of sequence-formation-edges = {null}
Pointer to next-pattern instance = {...}, Pointer to previous-pattern instance = {...}
}
```

Footnote:

4. sequence id 1 in accompanying sequence listing

FIG. 9A

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Start of first while loop

Bptr=0;seq="";offset=0;loopcnt=0;ht={};mr=1

Inside the loop

Roffset = 18;

Condition true -&gt; Inside 'if'

Seq = acttgatcggtagctaga<sup>5</sup>

Bptr= 18

Outside 'if'

poffset = 28

seq= acttgatcggtagctagacgcgcgcgcg<sup>6</sup>

ht={&lt;1,1&gt;}

loopcnt=1

mr=2

loopcnt=0

Start of second loop as mr!=null

Roffset = 28

Condition false

Poffset=39

Seq=acttgatcggtagctagacgcgcgcgcgaaataattaa<sup>7</sup>

ht={&lt;1,1&gt;,&lt;2,1&gt;}

loopcnt=1

mr=1

loopcnt=1

Start of third loop as mr!=null

Roffset = 39

Condition false

Poffset= 49

Seq= acttgatcggtagctagacgcgcgcgcgaaataattaaacgcgcgcgcg<sup>8</sup>

ht={&lt;1,2&gt;,&lt;2,1&gt;}

loopcnt=2

mr=3

loopcnt=0

Start of fourth loop as mr!=null

Roffset = 49

Condition false

Poffset=65

Seq= acttgatcggtagctagacgcgcgcgcgaaataattaaacgcgcgcgcgacaggtataggccaac<sup>9</sup>

ht={&lt;1,2&gt;,&lt;2,1&gt;,&lt;3,1&gt;}

loopcnt=1

mr=1

loopcnt=2

## Footnotes:

5. sequence id 2 in accompanying sequence listing
6. sequence id 3 in accompanying sequence listing
7. sequence id 4 in accompanying sequence listing
8. sequence id 5 in accompanying sequence listing
9. sequence id 6 in accompanying sequence listing

FIG. 9B

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Start of fifth loop as mr!=null

Roffset = 83

Condition true -> Inside 'if'

Seq=

acttgatcggtagctagacgcgcgcgcgaaataattaaacgcgcgcgcgacaggtataggccaaccggagaagctcccaaac<sup>10</sup>

Bptr=36

Outside 'if'

Poffset=93

Seq=

acttgatcggtagctagacgcgcgcgcgaaataattaaacgcgcgcgcgacaggtataggccaaccggagaagctcccaaacgcgcgcgcg<sup>11</sup>

ht={<1,3>,<2,1>,<3,1>}

loopcnt=3

mr=4

loopcnt=0

Start of sixth loop as mr!=null

Roffset =93

Condition false

Poffset=93

Seq=

acttgatcggtagctagacgcgcgcgcgaaataattaaacgcgcgcgcgacaggtataggccaaccggagaagctcccaaacgcgcgcgcgtactatatcatattac<sup>12</sup>

ht={<1,3>,<2,1>,<3,1>,<4,1>}

loopcnt=1

mr=null

loopcnt=-1

The while loop is terminated as mr = null;

Outside while loop

There is no more subsequence of the backbone to be added to 'Seq'

Return seq

Output =

"acttgatcggtagctagacgcgcgcgcgaaataattaaacgcgcgcgcgacaggtataggccaaccggagaagctcccaaacgcgcgcgcgtactatatcatattac"<sup>12</sup>

Footnotes:

10. sequence id 7 in accompanying sequence listing

11. sequence id 8 in accompanying sequence listing

12. sequence id 9 in accompanying sequence listing

FIG. 9C